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EXAMINER'S AMENDMENT

An examiner's amendment to the record appears below. Should the changes and/or additions be unacceptable to applicant, an amendment may be filed as provided by 37 CFR 1.312. To ensure consideration of such an amendment, it MUST be submitted no later than the payment of the issue fee. Authorization for this examiner's amendment was given in a telephone interview with James Voeller on1/28/10.

The application has been amended as follows:

44. (currently amended) Mutated A mutated V1/AR1/AV1 or C1/AL1/AC1 gene polynucleotide sequence of a tomato infecting geminivirus, wherein the mutations consist of silent point mutations distributed along the C1/AL1/AC1 polynucleotide sequence in such a way that the continuous homology between the mutated C1/AL1/AC1 polynucleotide sequence and the corresponding viral gene the wild-type C1/AL1/AC1 polynucleotide sequence is below less than or equal to 8 nucleotides, preferably below or equal to 5 nucleotides, said mutated C1/AL1/AC1 polynucleotide sequence encoding for a capsid protein or for a truncated Rep protein, respectively.

45-46. (canceled)

47. (currently amended) Mutated The mutated C1/AL1/AC1 gene polynucleotide sequence according to claim [[46]] 44, wherein the encoding truncated

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Rep proteins consisting of 130 amino acids (Rep_130) to 210 amino acids (Rep_210), said truncation occurring at the 3' terminal.

48. (currently amended) Mutated The mutated C1/AL1/AC1 gene polynucleotide sequence according to claim [[46]] 47, having the sequence of SEQ ID NO: 4, said polynucleotide sequence encoding for Rep-210 protein having the amino acid sequence of SEQ ID No 3 or SEQ ID No 5 NO: 5.

49. (cancelled)

- 50. (currently amended) Mutated The mutated gene C1/AL1/AC1 polynucleotide sequence according to claim 44 wherein the tomato infecting geminivirus is TYLCSV Tomato yellow leaf curl Sardinia virus (TYLCSV).
- **51. (currently amended)** Synthetic A construct comprising an heterologous polynucleotide sequence containing in the 5'-3' direction:
- a) <u>a</u> polynucleotide sequence acting as promoter in said plant, or <u>plant</u> tissue or transformed plant cells;
- b) a non_translated polynucleotide sequence positioned 5' of the encoding region of the mutated geminivirus C1/AL1/AC1 polynucleotide sequence;
- c) a mutated gene C1/AL1/AC1 polynucleotide sequence as defined according to claim 44; and

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d) a <u>polynucleotide</u> sequence acting as transcription terminator, positioned 3' with respect to the mutated <u>gene C1/AL1/AC1 polynucleotide sequence</u>.

- **52.** (currently amended) Expression An expression vector comprising the construct as defined according to claim 51.
- **53.** (currently amended) Transgenic A transgenic plant, or plant tissue or plant cells thereof, comprising in their genome a mutated gene C1/AL1/AC1 polynucleotide sequence as defined according to claim 44.
- **54.** (currently amended) Seed A seed comprising in its genome a mutated gene C1/AL1/AC1 polynucleotide sequence as defined according to claim 44.
- **70.** (New) The mutated C1/AL1/AC1 polynucleotide sequence according to claim 44, wherein the continuous homology between the mutated C1/AL1/AC1 polynucleotide sequence and the wild-type C1/AL1/AC1 polynucleotide sequence is less than or equal to 5 nucleotides.

Status of Rejections

All the rejections are withdrawn in light of the claims amendments and this amendment.

Allowable Subject Matter

Claims 44, 47-48, 50-55, 58-59, 63-65 and 68-70 are allowed.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Li Zheng whose telephone number is 571-272-8031. The examiner can normally be reached on Monday through Friday 9:00 AM - 5:30 PM EST.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Anne Marie Grunberg can be reached on 571-272-0975. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

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/Anne R Kubelik/ Primary Examiner, Art Unit 1638